#### Amendments to the Claims

This listing of the claims will replace all prior versions, and listings, of the claims in the application.

## Listing of Claims

### 1.-64. (cancelled)

- 65. (previously presented) A method for treating a lectin-mediated platelet disorder in a mammal comprising administering to the mammal a pharmaceutically effective amount of a nucleic acid ligand to P-selectin.
- 66. (currently amended) The method of claim 65 wherein said nucleic acid ligand to a leetin P-selectin is identified according to a method comprising:
- a) contacting a candidate mixture of nucleic acids with a lectin, wherein nucleic acids having an increased affinity to said lectin relative to the candidate mixture may be partitioned from the remainder of the candidate mixture;
- partitioning the increased affinity nucleic acids from the remainder of the candidate mixture; and
- c) amplifying the increased affinity nucleic acids to yield a mixture of nucleic acids enriched for nucleic acid sequences with relatively higher affinity and specificity for binding said lectin, whereby nucleic acid ligands of said lectin P-selectin may be identified.

#### 67. (cancelled)

 (currently amended) The method of claim 67 65 wherein said nucleic acid ligand to a leetin P-selectin is SEQ ID NO: 206.

- 69. (currently amended) A method for treating a lectin-mediated inflammation or lymphocyte tracking trafficking disorder in a mammal comprising administering to the mammal a pharmaceutically effective amount of a nucleic acid ligand to L-selectin.
- (currently amended) The method of claim 69 wherein said nucleic acid ligand to a leetin L-selectin is identified according to a method comprising:
- a) contacting a candidate mixture of nucleic acids with a lectin, wherein nucleic acids having an increased affinity to said lectin relative to the candidate mixture may be partitioned from the remainder of the candidate mixture;
- b) partitioning the increased affinity nucleic acids from the remainder of the candidate mixture; and
- c) amplifying the increased affinity nucleic acids to yield a mixture of nucleic acids enriched for nucleic acid sequences with relatively higher affinity and specificity for binding said lectin, whereby nucleic acid ligands of said lectin L-selectin may be identified.

# 71. (cancelled)

- 72. (currently amended) The method of claim 74 69 wherein said nucleic acid ligand to a-lectin L-selectin is SEQ ID NO: 185.
- 73. (new) The method of claim 68 wherein said nucleic acid ligand toP-selectin is a fragment of SEQ ID NO: 206 comprising a 38mer with a 5' end at position 19 and a 3' end at position 56.
- 74. (new) The method of claim 73 wherein said 38mer is further modified such that at least one of the guanines within said 38mer is 2'-O-methyl and at least one of said adenines within said 38mer is 2'-O-methyl.
- 75. (new) The method of claim 73 wherein said 38mer is further modified with 2'-OMe purine substitutions wherein said 38mer comprises:

# 5'-CUCAAC<u>GAG</u>CCAGG<u>A</u>ACAUCGACGUC<u>AGCA</u>AACGCGAG-3' (SEQ ID NO: 391)

wherein at least the underlined bases are 2'-OMe.

- 76. (new) A method for treating a lectin-mediated platelet disorder in a mammal comprising administering to said mammal a pharmaceutically effective amount of a formulation comprising a nucleic acid ligand to P-selectin wherein said ligand is a functional antagonist of PS-Rg.
- 77. (new) The method of claim 76 wherein said nucleic acid ligand to P-selectin is identified according to a method comprising:
- a) contacting a candidate mixture of nucleic acids with a lectin, wherein nucleic acids having an increased affinity to said lectin relative to the candidate mixture may be partitioned from the remainder of the candidate mixture;
- b) partitioning the increased affinity nucleic acids from the remainder of the candidate mixture: and
- c) amplifying the increased affinity nucleic acids to yield a mixture of nucleic acids enriched for nucleic acid sequences with relatively higher affinity and specificity for binding said lectin, whereby nucleic acid ligands of said P-selectin may be identified.
- $78. \qquad (\text{new}) \quad \text{The method of claim 76 wherein said nucleic acid ligand to P-selectin is} \\ \text{SEQ ID NO: } 206.$
- 79. (new) The method of claim 78 wherein said nucleic acid ligand to
  P-selectin is a fragment of SEQ ID NO: 206 comprising a 38mer with a 5'-end at position 19 and a 3'-end at position 56.
- 80. (new) The method of claim 79 wherein said 38mer is further modified such that at least one of the guanines within said 38mer is 2'-O-methyl and at least one of said adenines within said 38mer is 2'-O-methyl.

(new) The method of claim 79 wherein said 38mer is further modified with 2' OMe purine substitutions wherein said 38mer comprises;

5'-CUCAAC<u>GAG</u>CCAGG<u>A</u>ACAUCGACGUC<u>AG</u>C<u>A</u>AACGCGAG-3' (SEQ ID NO: 391)

wherein at least the underlined bases are 2'-OMe.

- 82. (new) A method for inhibiting the adhesion of platelets to neutrophils in the blood of a mammal comprising contacting a formulation comprising a nucleic acid ligand to Pselectin, wherein said ligand is a functional antagonist of PS-Rg, to said blood under conditions such that adherence of said platelets to said neutrophils is inhibited.
- 83. (new) The method of claim 82 wherein said nucleic acid ligand to P-selectin is identified according to a method comprising:
- a) contacting a candidate mixture of nucleic acids with a lectin, wherein nucleic acids having an increased affinity to said lectin relative to the candidate mixture may be partitioned from the remainder of the candidate mixture;
- b) partitioning the increased affinity nucleic acids from the remainder of the candidate mixture; and
- c) amplifying the increased affinity nucleic acids to yield a mixture of nucleic acids enriched for nucleic acid sequences with relatively higher affinity and specificity for binding said lectin, whereby nucleic acid ligands of said P-selectin may be identified.
- $84. \hspace{0.5cm}$  (new) The method of claim 82 wherein said nucleic acid ligand to P-selectin is SEQ ID NO: 206.
- 85. (new) The method of claim 84 wherein said nucleic acid ligand to
  P-selectin is a fragment of SEQ ID NO: 206 comprising a 38mer with a 5'-end at position 19 and a 3'-end at position 56.

86. (new) The method of claim 85 wherein said 38mer is further modified such that at least one of the guanines within said 38mer is 2'-O-methyl and at least one of said adenines within said 38mer is 2'-O-methyl.

87. (new) The method of claim 85 wherein said 38mer is further modified with 2'-OMe purine substitutions wherein said 38mer comprises:

5'-CUCAAC<u>GAG</u>CCAGG<u>A</u>ACAUCGACGUC<u>AG</u>C<u>A</u>AACGCGAG-3' (SEQ ID NO: 391)

wherein at least the underlined bases are 2'-OMe.

- 88. (new) A method for inhibiting the adhesion of platelets to leukocytes in the blood of a mammal comprising administering to said mammal a nucleic acid ligand to P-selectin.
- 89. A method for inhibiting the binding of P-selectin to a carbohydrate in the blood of a mammal comprising administering to said mammal a nucleic acid ligand to P-selectin.